SCORE Search Results Details for Application 10552515 and Search Result 20090316_112342_us-10-552-515-2.rni.

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This page gives you Search Results detail for the Application 10552515 and Search Result 20090316_112342_us-10-552-515-2. rni.

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OM nucleic - nucleic search, using sw model

Run on: March 16, 2009, 16:27:41; Search time 1151 Seconds

(without alignments)

20454.958 Million cell updates/sec

Title: US-10-552-515-2

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Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 9875436 seqs, 3558593875 residues

Total number of hits satisfying chosen parameters: 19750872

Minimum DB seg length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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	2	461	13.9	13243	8	US-10-741-600-17879	Sequence 17879, A
	3	461	13.9	14172	6	US-10-741-601-5626	Sequence 5626, Ap
	4	461	13.9	14172	8	US-10-741-600-17603	Sequence 17603, A
С	5	460.6	13.9	101046	6	US-10-741-601-5689	Sequence 5689, Ap
С	6	460.6	13.9	101046	8	US-10-741-600-17753	Sequence 17753, A
	7	432.8	13.1	4509	8	US-10-912-745B-698	Sequence 698, App
	8	325.6	9.8	3052	5	US-10-342-887-1730	Sequence 1730, Ap
	9	301.6	9.1	3898	3	US-10-104-047-604	Sequence 604, App
	10	286.6	8.7	2736	3	US-10-104-047-571	Sequence 571, App
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	13	216.2	6.5	1282	3	US-09-270-767-13982	Sequence 13982, A
	14	170.8	5.2	2293	3	US-10-104-047-1146	Sequence 1146, Ap
	15	157.2	4.8	2371	7	US-10-100-683-1599	Sequence 1599, Ap
	16	157.2	4.8	2371	7	US-11-001-793-1599	Sequence 1599, Ap
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	37	58.2	1.8	7218	2	US-08-232-463-14	Sequence 14, Appl
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ALIGNMENTS

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; Sequence 5735, Application US/10741601
; Patent No. 7306913
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
 TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
 TITLE OF INVENTION: STENOSIS, METHODS OF DETECTION AND USES THEREOF
  FILE REFERENCE: CL001500
; CURRENT APPLICATION NUMBER: US/10/741,601
  CURRENT FILING DATE: 2003-12-22
; NUMBER OF SEQ ID NOS: 26415
  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 5735
  LENGTH: 13243
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  ORGANISM: Homo sapiens
  FEATURE:
  NAME/KEY: misc feature
  LOCATION: (1)...(13243)
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; Sequence 17879, Application US/10741600
; Patent No. 7482117
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
  TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
; TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF DETECTION AND USES THEREOF
 FILE REFERENCE: CL001499
; CURRENT APPLICATION NUMBER: US/10/741,600
; CURRENT FILING DATE: 2003-12-22
; NUMBER OF SEO ID NOS: 73997
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 17879
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  FEATURE:
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  LOCATION: (1)...(13243)
; OTHER INFORMATION: n = A.T.C or G. or insertion/deletion polymorphism (see Tables 1-2)
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; Sequence 5626, Application US/10741601
; Patent No. 7306913
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
 TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
 TITLE OF INVENTION: STENOSIS, METHODS OF DETECTION AND USES THEREOF
 FILE REFERENCE: CL001500
; CURRENT APPLICATION NUMBER: US/10/741,601
; CURRENT FILING DATE: 2003-12-22
; NUMBER OF SEQ ID NOS: 26415
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; SEO ID NO 5626
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; Sequence 17603, Application US/10741600
; Patent No. 7482117
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
 TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
 TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001499
; CURRENT APPLICATION NUMBER: US/10/741,600
; CURRENT FILING DATE: 2003-12-22
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; Patent No. 7306913
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
 TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
 TITLE OF INVENTION: STENOSIS, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001500
 CURRENT APPLICATION NUMBER: US/10/741,601
; CURRENT FILING DATE: 2003-12-22
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  TYPE: DNA
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; Sequence 17753, Application US/10741600
; Patent No. 7482117
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
: TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
 TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001499
; CURRENT APPLICATION NUMBER: US/10/741,600
; CURRENT FILING DATE: 2003-12-22
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; SEO ID NO 17753
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  OTHER INFORMATION: n = A,T,C or G, or insertion/deletion polymorphism (see Tables 1-2)
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US-10-912-745B-698
; Sequence 698, Application US/10912745B
; Patent No. 7473531
; GENERAL INFORMATION
; APPLICANT: DOMON, Bruno et al.
; TITLE OF INVENTION: Pancreatic Cancer Targets and Uses
; TITLE OF INVENTION: Thereof
; FILE REFERENCE: CL001538
; CURRENT APPLICATION NUMBER: US/10/912,745B
; CURRENT FILING DATE: 2004-08-06
: NUMBER OF SEO ID NOS: 875
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RESHLT 8

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; Sequence 1730, Application US/10342887
; Patent No. 7171311
; GENERAL INFORMATION:
; APPLICANT: Dai, Hongyue
; APPLICANT: He, Yudong
; APPLICANT: Linsley, Peter S.
; APPLICANT: Mao, Mao
; APPLICANT: Roberts, Christopher J.
; APPLICANT: Van 't Veer, Laura Johanna
; APPLICANT: Van de Vijver, Marc J.
; APPLICANT: Bernards, Rene
; TITLE OF INVENTION: Diagnosis and Prognosis of Breast Cancer Patients
; FILE REFERENCE: 9301-188-999
; CURRENT APPLICATION NUMBER: US/10/342,887
; CURRENT FILING DATE: 2003-01-15
; PRIOR APPLICATION NUMBER: 60/298,918
; PRIOR FILING DATE: 2001-06-18
; PRIOR APPLICATION NUMBER: 60/380,710
; PRIOR FILING DATE: 2002-05-14
; PRIOR APPLICATION NUMBER: 10/172,118
; PRIOR FILING DATE: 2002-06-14
; NUMBER OF SEQ ID NOS: 2699
; SEO ID NO 1730
; LENGTH: 3052
; TYPE: DNA
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 Matches 759; Conservative 0; Mismatches 589; Indels 30; Gaps 5;
Qv
      1235 AGACATACCCACGCAGGAACTGTGTGGCAGCAAGGACAGCTTCGAGATGTGCCCACTTTG 1294
          Db
         7 AAACATCCCCAGCATGGAGATGTGTGACCAGAGACACAATATCACCATGTGCCCGCTTTG 66
Qv
      Db
        1352 GCTGTTCGACCACGGCGCACCGTGTTCTTCAGCTTGTTCATGGCACTGTGGGCCGTGCT 1411
Qу
           Db
       127 CCTCTTCGACACCCCGCCACGGTCTTCTTCTCTGTCTTCATGGCCCTCTGGGCTGCCAC 186
      1412 GCTGCTGGAGTACTGGAAGCGGAAGAGCGCCACGCTGGCCTACCGCTGGGACTGCTCTGA 1471
0v
            Db
      187 CTTCATGGAGCACTGGAAGCGGAAACAGATGCGACTCAACTACCGCTGGGACCTCACGGG 246
    1472 CTACGAGGACACTGAGGAG------AGGCCTCGGCCCCAGTTTGCCGCCTCAGC 1519
Qv
          THE THE THEFT
                                    247 CTTTGAAGAGGAAGAGGAGCTGTCAAGGATCATCCTAGAGCTGAATACGAAGCCAGAGT 306
Db
```

Qy	1520	CCCCATGACAGCCCCGAACCCCATCACGGGTGAGGACGAGCCCTACTTCCCTGAGAGGAG	1579
Db	307		366
Qy	1580	CCGCGCGCGCCGCATGCTGGCCGGCTCTGTGGTGATCGTGGTGATGGTGGC	1630
Db	367	ATGGAGAGATCGGTTCCCAGCCTACCTCACTACTTGGTCTCCATCATCTTCATGATTGC	426
QУ	1631	CGTGGTGGTCATGTGCCTCGTGTCTATCATCCTGTACCGTGCCATCATGGCCATCGTGGT	1690
Db	427	AGTGACGTTTGCCATCGTCCTCGGCGTCATCATCTACAGGATCTCCATGGCCGCCGCCTT	486
Qy	1691	GTCCAGGTCGGGCAACACCCTTCTCGCAGCCTGGGCCTCTCGCATCGCCAGCCTCACGGG	1750
Db	487	GGCCATGAACTCCTCCCCTCCGTGCGGTCCAACATCCGGGTCACAGTCACAGCCACCGC	546
Qy	1751	GTCTGTAGTGAACCTCGTCTTCATCCTCATCCTCCAAGATCTATGTATCCCTGGCCCA	1810
Db	547	${\tt GGTCATCATCAACCTAGTGGTCATCATCCTCCTGGACGAGGTGTATGGCTGCATAGCCCG}$	606
Qy	1811	CGTCCTGACACGATGGGAAATGCACCGCACCCAGACCAAGTTCGAGGACGCCTTCACCCT	1870
Db	607	$\tt ATGGCTCACCAAGATCGAGGTCCCAAAGACGGAGAAAAGCTTTGAGGAGAGGGCTGATCTT$	666
Qy	1871	CAAGGTGTTCATCTTCCAGTTCGTCAACTTCTACTCCTCACCCGTCTACATTGCCTTCTT	1930
Db	667	${\tt CAAGGCTTTCCTGCTGAAGTTTGTGAATTCCTACACCCCCATCTTTTACGTGGCGTTCTT}$	726
Qу	1931	CAAGGGCAGGTTTGTGGGATACCCAGGCAACTACCACACCTTGTTTGGAGTCCGCAA	1987
Db	727	$\tt CAAAGGCCGGTTTGTTGGACGCCCGGGCGACTACGTGTACATTTTCCGTTCCTTCC$	786
Qy	1988	TGAGGAGTGCGCGGCTGGAGCTGCCTGATCGAGCTGGCACAGGAGCTCCTGGTCATCAT	2047
Db	787	GGAAGAGTGTGCGCCAGGGGGCTGCCTGATGGAGCTATGCATCCAGCTCAGCATCATCAT	
Qy	2048	GGTGGGCAAGCAGTCATCAACAACATGCAGGAGGTCCTCATCCCGAAGCTAAAGGG	2104
Db	847	GCTGGGGAAACAGCTGATCCAGAACAACCTGTTCGAGATCGGCATCCCGAAGATGAAGAA	906
Qy	2105	CTGGTGGCAGAAGTTCCGGCTTCGCTCCAAGAAGAGGAGGCGGGAGCTTCTGCAGGGGC	2164
Db	907	$\tt GCTCATCCGCTACCTGAAGCTGAAGCAGCAGGAGCCCCCCTGACCACGAGGAGTGTGTGAA$	966
Qy	2165	TAGCCAGGGGCCCTGGGAGGACGACTATGAGCTTGTGCCCTGTGAGGGTCTGTTTGACGA	2224
Db	967	GAGGAAACAGCGGTACGAGGTGGATTACAACCTGGAGCCCTTCGCGGGCCTCACCCCAGA	1026
Qy	2225	GTACCTGGAAATGGTGCTGCAGTTCGGCTTCGTCACCATCTTCGTGGCCGCCTGTCCGCT	2284
Db	1027	${\tt GTACATGGAAATGATCATCCAGTTTGGCTTCGTCACCCTGTTTGTCGCCTCCTTCCCCCT}$	1086
Qy	2285	CGCGCCGCTCTTCGCCCTGCTCAACAACTGGGTGGAGATCCGCTTGGACGCGCGCAAGTT	2344
Db	1087	$\tt GGCCCCACTGTTTGCGCTGCACAACAACATCATCGAGATCCGCCTGGACGCCAAAAAGTT$	1146

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2345 CGTCTGCGAGTACCGGCGCCCTGTGGCCGAGCGCCCCAGGACATCGGCATCTGGTTCCA 2404
Qv
               Db
      1147 TGTCACTGAGCTCCGAAGGCCGGTAGCTGTCAGAGCCAAAGACATCGGAATCTGGTACAA 1206
      2405 CATCCTGGCGGGCCTCACGCACCTGGCGGTCATCAGCAACGCCTTCCTCCTGGCCTTCTC 2464
Qy
           Db
      1207 TATCCTCAGAGGCATTGGGAAGCTTGCTGTCATCATCAATGCCTTCGTGATCTCCTTCAC 1266
      2465 GTCCGACTTCCTGCCGCGCCCTACTACCGGTGGACCCGCGCCCACGACCTGCGCGGCTT 2524
QУ
          Db
      1267 GTCTGACTTCATCCCGCGCCTGGTGTACCTCTACATGTACAGTAAGAACGGGACCATGCA 1326
      Qv
          Db
      1327 CGGCTTCGTCAACCACCCCTCTCCTCCTTCAACGTCAGTGACTTCCAGAACGGCACG 1384
RESULT 9
US-10-104-047-604
; Sequence 604, Application US/10104047
; Patent No. 6943241
: GENERAL INFORMATION:
: APPLICANT: HELIX RESEARCH INSTITUTE
 TITLE OF INVENTION: No. 6943241el full length cDNA
; FILE REFERENCE: H1-A0105
 CURRENT APPLICATION NUMBER: US/10/104,047
; CURRENT FILING DATE: 2002-03-25
; PRIOR APPLICATION NUMBER:
; PRIOR FILING DATE:
; NUMBER OF SEO ID NOS: 4096
; SOFTWARE: PatentIn Ver. 2.1
: SEO ID NO 604
; LENGTH: 3898
; TYPE: DNA
 ORGANISM: Homo sapiens
US-10-104-047-604
 Ouerv Match
                   9.1%; Score 301.6; DB 3; Length 3898;
 Best Local Similarity 50.9%; Pred. No. 8e-54;
 Matches 871; Conservative 0; Mismatches 824; Indels 15; Gaps 6;
       780 GAGTACTACTCCTGCCGGTTCAGAGTGAACAAGCTGCCACGCTTCCTCGGGAGTGACAAC 839
Qу
          Db
       950 GACTGCTACACTGCCCCTTTCAGCCAGCAAAGGATCCATCACTTCATC---ATACACAAC 1006
       840 CAGGACACCTTCTTCACAAGCACCAAGAGGCACCAAATTCTGTTTGAGATCCTGGCCAAG 899
0v
           Db
      1007 AAAGAAACGTTCTTCAACAATGCCACAAGAAGTAGAATCGTGCATCACATTTTACAAAGA 1066
      900 ACCCCGTATGGCCACGAGAAGAAAACCTGCTTGGGATCCACCAGCTGCTGGCAGAGGGT 959
Qv
               Db
      1067 ATAAAATATG---AAGAAGGAAAAAACAAGATTGGTCTGAATCGTTTGCTTACCAATGGC 1123
       960 GTCCTCAGTGCCGCCTTCCCCTGCATGACGGCCCCTTCAAGACGCCCCCAGAGGGCCCG 1019
Ov
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1904 ACCACAGGGACTGCTGTGCCATCAACTTCTGTATCATTATGTTGCTGAATGTGCTCTAT 1963 1797 GTATCCCTGGCCCACGTCCTGACACGATGGGAAATGCACCGCACCCAGACCAAGTTCGAG 1856

11 1

Qу

Db

Ov

```
Db
      1964 GAAAAAGTTGCCCTGCTTCTGACGAATTTAGAACAGCCTCGCACAGAGTCTGAGTGGGAG 2023
      1857 GACGCCTTCACCCTCAAGGTGTTCATCTTCCAGTTCGTCAACTTCTACTCCTCACCCGTC 1916
Qy
           2024 AACAGCTTCACCCTGAAAATGTTTCTTTTTCAGTTTGTCAATCTGAACAGCTCCACATTT 2083
Db
      1917 TACATTGCCTTCTTCAAGGGCAGGTTTGTGGGATACCCAGGCAACTACCACACCTTGTT- 1975
          Db
      2084 TACATCGCATTCTTCCTCGGAAGATTTACAGGACACCCAGGTGCCTACTTGAGGCTGATA 2143
      1976 --TGGAGTCCGCAATGAGGAGTGCGCGGCTGGAGGCTGCCTGATCGAGCTGGCACAGGAG 2033
Qv
             1
                - 1
                    2144 AACAGGTGGAGACTAGAAGAGTGCCACCCTAGTGGATGCCTTATTGATCTGTGTATGCAA 2203
Db
      2034 CTCCTGGTCATCATGGTGGGCAAGCAGGTCATCAACAACATGCAGGAGGTCCTCATCCCG 2093
Qv
               2204 ATGGGTATTATAATGGTGCTAAAGCAGACCTGGAATAATTTCATGGAACTTGGCTACCCG 2263
Db
      2094 AAGCTAAAGGGCTGGTGGCAGAAGTTCCGGCTTCGCTCCAAGAAGAGGAAGGCGGGAGCT 2153
0v
             Db
      2264 TTAATTCAGAATTGGTGGACTAGAAGAAAGTACG--ACAAGAACATGGACCTGAAAGGA 2321
Qу
      2154 TCTGCAGGGGCTAGCCAGGGGCCCTGGGAGGACGACTATGAGCTTGTGCCCTGTGAGGGT 2213
              Dh
      2322 AAATAAGTTTCCCACAATGGGAAA-AGGACTATAACCTTCAGCCGATGAATGCCTATGGA 2380
Qv
      2214 CTGTTTGACGAGTACCTGGAAATGGTGCTGCAGTTCGGCTTCGTCACCATCTTCGTGGCC 2273
          2381 CTCTTCGATGAATACTTAGAAATGATTCTTCAGTTTGGATTCACAACTATCTTTGTGGCA 2440
Db
Qу
      2274 GCCTGTCCGCTCGCCGCTCTTCGCCCTGCTCAACAACTGGGTGGAGATCCGCTTGGAC 2333
          2441 GCTTTTCCCCTAGCACCACTTCTGGCCTTACTGAATAACATAATTGAAATTCGACTTGAT 2500
Db
      2334 GCGCGCAAGTTCGTCTGCGAGTACCGGCGCCCTGTGGCCGAGCGCGCCCAGGACATCGGC 2393
Qу
                            1.1
              Db
      2501 GCTTACAAATTTGTCACACAGTGGAGGAGACCTTTAGCTTCAAGGGCCAAAGACATAGGA 2560
      2394 ATCTGGTTCCACATCCTGGCGGGCCTCACGCACCTGGCGGTCATCAGCAACGCCTTCCTC 2453
Qv
          2561 ATTTGGTATGGAATTCTTGAAGGCATTGGAATTCTCTCTGTTATCACAAATGCATTTGTC 2620
Db
      2454 CTGGCCTTCTCGTCCGACTTCCTGCCGCGC 2483
Qу
           Db
      2621 ATAGCGATAACATCTGACTTTATCCCTCGC 2650
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RESULT 10
US-10-104-047-571
; Sequence 571, Application US/10104047
; Patent No. 6943241
; GENERAL INFORMATION:
  APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: No. 6943241el full length cDNA
  FILE REFERENCE: H1-A0105
  CURRENT APPLICATION NUMBER: US/10/104,047
```

; CURRENT FILING DATE: 2002-03-25 ; PRIOR APPLICATION NUMBER:

```
; PRIOR FILING DATE:
; NUMBER OF SEO ID NOS: 4096
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 571
 LENGTH: 2736
  TYPE: DNA
 ORGANISM: Homo sapiens
US-10-104-047-571
 Query Match
                   8.7%; Score 286.6; DB 3; Length 2736;
 Best Local Similarity 51.7%; Pred. No. 1.1e-50;
 Matches 752; Conservative 0; Mismatches 694; Indels 9; Gaps 4;
     1035 AACCAGCGCCAAGTCCTTTTCCAGCACTGGGCGCGCTGGGGCAAGTGGAACAAGTACCAG 1094
Qу
          Db
        31 AACCACCGACATCTACTCTATGAGTGCTGGGCCTCCTGGGGCGTGTGGTATAAATACCAA 90
      1095 CCCCTGGACCACGTGCGCAGGTACTTCGGGGAGAAGGTGGCCCTCTACTTCGCCTGGCTC 1154
Qv.
          91 CCTTTGGATCTTGTAAGGCGGTACTTTGGAGAGAAGATTGGGTTATATTTTGCCTGGTTG 150
Db
      1155 GGGTTTTACACAGGCTGGCCCGCCAGCGGCAGTGGTGGCACACTGGTGTTCCTGGTG 1214
Qy
          Db
      151 GGCTGGTACACCGGCATGCTCTTCCCAGCTGCCTTCATTGGATTGTTTTGTCTTTTTGTAT 210
      1215 GGCTGCTTCCTGGTGTTCTCAGACATACCCACGCAGGAACTGTGTGGCAGCAAGGACAGC 1274
Οv
          Db
       211 GGCGTCACCACTCTGGATCACAGCCAAGTCAGTAAAGAAGTCTGCCAAGCTACAGATATC 270
      1275 TTCGAGATGTGCCCACTTTGCCTCGACTGCCCTTTCTGGCTGCTCTCCAGCGCCTGTGCC 1334
Qv.
           Db
       271 ATCATGTGTCCTGTGTGTATAAATACTGTCCATTCATGAGGCTGTCAGACAGCTGTGTA 330
      1335 CTGGCCCAGGCCGGCCGCTGTTCGACCACGGCGGCACCGTGTTCTTCAGCTTGTTCATG 1394
Qv
            Db
       331 TATGCCAAGGTAACCCACCTTTTTGACAATGGAGCCACTGTCTTCTTTGCTGTTTTCATG 390
      1395 GCACTGTGGGCCGTGCTGCTGGAGTACTGGAAGCGGAAGAGCGCCACGCTGGCCTAC 1454
Οv
          Db
       391 GCAGTCTGGGCAACAGTTTTCCTGGAGTTTTGGAAAAGACGGCGAGCAGTAATTGCTTAT 450
      1455 CGCTGGGACTGCTCTGACTACGAGGACACTGAGGAGGGCCTCGGCCCCAGTTTGCCGCC 1514
0v
            451 GACTGGGATTTGATAGACTGGGAAGAAGAGGAGGAGAAATACGACCCCAGTTTGAAGCC 510
Db
      1515 TCAGCCCCCATGACAGCCCC---GAACCCCATCACGGGTGAGGACGAGCCCTACTTCCCT 1571
Qv
                511 AAGTATTCCAAGAAAGAGCGGATGAATCCAATTTCTGGAAAGCCAGAACCTTATCAAGCA 570
Db
Qy
      1572 GAGAGGAGCCGCGCGCGCCCCATGCTGGCCGGCTCTGTGGTGATCGTGGTGATGGTGGCC 1631
                      Db
      571 TTTACAGATAAATGCAGCAGACTTATCGTTTCTGCATCTGGAATATTTTTTATGATCTGC 630
Ov
      1632 GTGGTGGTCATGTGCCTCGTGTCTATCATCCTGTACCGTGCCATCATGGCCATCGTGGTG 1691
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Db	631	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	690
Qy	1692	TCCAGGTCGGGCAACACCCTTCTCGCAGCCTGGGCCTCTCGCATCGCCAGCCTCACGGGG	1751
Db	691	GCTGCCTTTAAGTGGGCGTTAATCAGGAATAACTCTCAGGTTGCAACCACAGGGACTGCT	750
Qу	1752	TCTGTAGTGAACCTCGTCTTCATCCTCTCCCAAGATCTATGTATCCCTGGCCCAC	1811
Db	751	GTGTGCATCAACTTCTGTATCATTATGTTGCTGAATGTGCTCTATGAAAAAGTTGCCCTG	810
Qy	1812	GTCCTGACACGATGGGAAATGCACCGCACCCAGACCAAGTTCGAGGACCCTTCACCCTC	1871
Db	811	CTTCTGACGAATTTAGAACAGCCTCGCACAGAGTCTGAGTGGGAGAACAGCTTCACCCTG	
Qy	1872	AAGGTGTTCATCTTCCAGTTCGTCAACTTCTACTCCTCACCCGTCTACATTGCCTTCTTC	1931
Db	871	AAAATGTTTCTTTTCAGTTTGTCAATCTGAACAGCTCCACATTTTACATCGCATTCTTC	
Qy	1932	AAGGGCAGGTTTGTGGGATACCCAGGCAACTACCACCTTGTTTGGAGTCCGCAAT	1988
Db	931	CTCGGAAGATTTACAGGACACCCAGGTGCCTACTTGAGGCTGATAAACAGGTGGAGACTA	
Qy	1989	GAGGAGTGCGCGGCTGGAGCTGCTGATCGAGCTGGCACAGGAGCTCCTGGTCATCATG	2048
Db	991	${\tt GAAGAGTGCCACCCTAGTGGATGCCTTATTGATCTGTGTATGCAAATGGGTATTATAATG}$	1050
Qy	2049	GTGGGCAAGCAGGTCATCAACAACATGCAGGAGGTCCTCATCCCGAAGCTAAAGGGCTGG	2108
Db	1051	$\tt GTGCTAAAGCAGACCTGGAATAATTCATGGAACTTGGCTACCCGTTAATTCAGAATTGG$	1110
Qy	2109	$ \begin{tabular}{lllllllllllllllllllllllllllllllllll$	2168
Db	1111	${\tt TGGACTAGAAGAAAGTACGACAAGAACATGGACCTGAAAGGAAAATAAGTTTCCCAC}$	1168
QУ	2169	CAGGGGCCCTGGGAGGACGACTATGAGCTTGTGCCCTGTGAGGGTCTGTTTGACGAGTAC	2228
Db	1169	${\tt AATGGG-AAAAGGACTATAACCTTCAGCCGATGAATGCCTATGGACTCTTCGATGAATAC}$	1227
Qу	2229	CTGGAAATGTGCTGCAGTTCGGCTTCGTCACCATCTTCGTGGCCGCCTGTCCGCC	2288
Db	1228	${\tt TTAGAAATGATTCTTCAGTTTGGATTCACAACTATCTTTGTGGCAGCTTTTCCCCTAGCA}$	1287
Qy	2289	CCGCTCTTCGCCCTGCTCAACAACTGGGTGGAGATCCGCTTGGACGCGCGCAAGTTCGTC	2348
Db	1288	CCACTTCTGGCCTTACTGAATAACATAATTGAAATTCGACTTGATGCTTACAAATTTGTC	
Qу	2349	TGCGAGTACCGGCGCCCTGTGGCCGAGCGCCCAGGACATCGGCATCTGGTTCCACATC	2408
Db	1348	${\tt ACACAGTGGAGGAGACCTTTAGCTTCAAGGGCCAAAGACATAGGAATTTGGTATGGAATT}$	1407
Qу	2409	CTGGCGGGCCTCACGCACCTGGCGTCATCAGCAACGCCTTCCTCCTGGCCTTCTCGTCC	2468
Db	1408	$\tt CTTGAAGGCATTGGAATTCTCTCTGTTATCACAAATGCATTTGTCATAGCGATAACATCT$	1467
Qy	2469	GACTTCCTGCCGCGC 2483	

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RESULT 11
US-10-108-260A-2040
; Sequence 2040, Application US/10108260A
; Patent No. 7193069
; GENERAL INFORMATION:
; APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: No. 7193069el full length cDNA
; FILE REFERENCE: H1-A0106
; CURRENT APPLICATION NUMBER: US/10/108,260A
; CURRENT FILING DATE: 2002-03-27
; NUMBER OF SEQ ID NOS: 5458
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2040
 LENGTH: 2118
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-108-260A-2040
 Query Match
                   7.6%; Score 252.6; DB 5; Length 2118;
 Best Local Similarity 54.3%; Pred. No. 1.7e-43;
 Matches 616; Conservative 0; Mismatches 489; Indels 30; Gaps 4;
Qv
      841 AGGACACCTTCTCACAAGCACCAAGAGGCACCAAATTCTGTTTGAGATCCTGGCCAAGA 900
          Db
       731 AGGATTCCTTTTCGACAGCAAAACCCGGAGCACGATTGTCTATGAGATCTTGAAGAGAA 790
      901 CCCCGTATGGCCACGAGAAGAAAACCTGCTTGGGATCCACCAGCTGCTGGCAGAGGGTG 960
Qу
          Db
      791 CGACGTGTACAAAGGCCAAGTACAGCATG---GGCATCACGAGCCTGCTGGCCAATGGTG 847
      961 TCCTCAGTGCCGCCTTCCCCTGCATGACGCCCCTTCAAGACGCCCCCAGAGGGCCCGC 1020
Qv
                Db
      848 TGTACGCGGCTGCATACCCACTGCACGATGGAGACTACAACGGTGAAAACGTCGAGT--- 904
     1021 AGGCTCCACGCCTCAACCAGCGCCAAGTCCTTTTCCAGCACTGGGCGCGCTGGGGCAAGT 1080
Qv
                   905 -----TCAACGACAGAAAACTCCTGTACGAAGAGTGGGCACGCTATGGAGTTT 952
Db
      1081 GGAACAAGTACCAGCCCTGGACCACGTGCGCAGGTACTTCGGGGAGAAGGTGGCCCTCT 1140
Qу
             Db
       953 TCTATAAGTACCAGCCCATCGACCTGGTCAGGAAGTATTTTGGGGAGAAGATCGGCCTGT 1012
      1141 ACTTCGCCTGGCTCGGGTTTTACACAGGCTGGCTCCTGCCAGCGGCAGTGGTGGGCACAC 1200
0v
          Db
      1013 ACTTCGCCTGGCTGGGCGTGTACACCCAGATGCTCATCCCTGCCTCCATCGTGGGAATCA 1072
     1201 TGGTGTTCCTGGTGGGCTGCTTCCTGGTGTTCTCAGACATACCCACGCAGGAACTGTGTG 1260
Qv
          Db
      1073 TTGTCTTCCTGTACGGATGCGCCACCATGGATGAAAACATCCCCAGCATGGAGATGTGTG 1132
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Ov

1261 GCAGCAAGGACAGCTTCGAGATGTGCCCACTTTGCCTCGA---CTGCCCTTTCTGGCTGC 1317

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Db
      1133 ACCAGAGACACAATATCACCATGTGCCCGCTTTGCGACAAGACCTGCAGCTACTGGAAGA 1192
      1318 TCTCCAGCGCCTGTGCCCTGGCCCAGGCCGGCTGTTCGACCACGGCGGCACCGTGT 1377
Qy
          1193 TGAGCTCAGCCTGCGCCACGGCCGCCACCTCTTCGACAACCCCGCCACGGTCT 1252
Db
      1378 TCTTCAGCTTGTTCATGGCACTGTGGGCCGTGCTGCTGCAGTACTGGAAGCGGAAGA 1437
QУ
          1253 TCTTCTCTGTCTTCATGGCCCTCTGGGCTGCCACCTTCATGGAGCACTGGAAGCGGAAAC 1312
      1438 GCGCCACGCTGGCCTACCGCTGGGACTGCTCTGACTACGAGGACACTGAGGAGAGGCCTC 1497
Qv
                1313 AGATGCGACTCAACTACCGCTGGGACCTCACGGGCTTTGAAGAGGAAGAGGATCATCCTA 1372
Db
      1498 GGCCCCAGTTTGCCGCCTCAGCCCCCATGACAGCCCCGAACCCCATCACGGGTGAGGACG 1557
Qv
          1373 GAGCTGAATACGAAGCCAGAGTCTTGGAGAAGTCTCTGAAGAAGAGTCCAGAAACAAAG 1432
Db
      1558 AGCCCTACTTCCCTGAGAGGAGCCGCGCGCCGCATGCTGGCC-----GGCTCTG 1608
Qу
          11 1
      1433 AGACTGACAAAGTGAAGCTGACATGGAGAGATCGGTTCCCAGCCTACCTCACTAACTTGG 1492
Db
Qу
      1609 TGGTGATCGTGGTGGTGGCCGTGGTGGTCATGTGCCTCGTGTCTATCATCCTGTACC 1668
             Db
      1493 TCTCCATCATCTTCATGATTGCAGTGACGTTTGCCATCGTCCTCGGCGTCATCATCTACA 1552
Qv
      1669 GTGCCATCATGGCCATCGTGGTGTCCAGGTCGGGCAACACCCTTCTCGCAGCCTGGGCCT 1728
           1553 GGATCTCCATGGCCGCCCCTTGGCCATGAACTCCTCCCCCTCCGTGCGGTCCAACATCC 1612
Db
Qу
      1729 CTCGCATCGCCAGCCTCACGGGGTCTGTAGTGAACCTCGTCTTCATCCTCATCCTCTCCA 1788
            1613 GGGTCACAGTCACAGCCACCGCGGTCATCATCAACCTAGTGGTCATCATCCTCCTGGACG 1672
Db
      1789 AGATCTATGTATCCCTGGCCCACGTCCTGACACGATGGGAAATGCACCGCACCCAGACCA 1848
Qу
          11 111 1
Db
      1673 AGGTGTATGGCTGCATAGCCCGATGGCTCACCAAGATCGAGGTCCCAAAGACGGAGAAAA 1732
      1849 AGTTCGAGGACGCCTTCACCCTCAAGGTGTTCATCTTCCAGTTCGTCAACTTCTACTCCT 1908
Qv
           1733 GCTTTGAGGAGAGGCTGATCTTCAAGGCTTTCCTGCTGAAGTTTGTGAATTCCTACACCC 1792
Db
      1909 CACCCGTCTACATTGCCTTCTTCAAGGGCAGGTTTGTGGGATACCCAGGCAACTA 1963
QУ
            Db
      1793 CCATCTTTTACGTGGCGTTCTTCAAAGGCCGGTTTGTTGGACGCCCGGGCGACTA 1847
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RESULT 12
US-10-108-260A-1547; Sequence 1547, Application US/10108260A; Patent No. 7193069; GENERAL INFORMATION:
APPLICANT: HELIX RESEARCH INSTITUTE; TITLE OF INVENTION: No. 7193069e1 full length cDNA; FILE REFERENCE: H1-A0106; CURENT APPLICATION NOMBER: US/10/108.260A
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CURRENT FILING DATE: 2002-03-27
; NUMBER OF SEQ ID NOS: 5458
; SOFTWARE: Patentin Ver. 2.1
; SEO ID NO 1547
 LENGTH: 2158
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-108-260A-1547
 Ouerv Match
                  7.2%; Score 239.2; DB 5; Length 2158;
 Best Local Similarity 52.3%; Pred. No. 1.2e-40;
 Matches 693; Conservative 0; Mismatches 568; Indels 63; Gaps 5;
     1508 TGCCGCCTCAGCCCCCATGACAGCCCCGAACCCCATCACGGGTGAGGACGAGCCCTACTT 1567
Qv
          276 TGCCGTGTCTGAGGAGGAAATGGCACTTCAGCTCATTAACTGCCCCGACTACAAGCTCCG 335
Db
      1568 CCCTGAGAGGAGCCGCGCGCGCGCATGCTGGCCGGCTCTGTGGTGATCGTGGTGATGGT 1627
Qv
           336 GCCATACCAGCACTCCTACCTACGCAGCACCGTCATCCTCGTCCTGACCCTGCTCATGAT 395
Db
     1628 GGCCGTGGTGGTCATGTGCCTCGTGTCTATCATCCTGTACCGTGCCATCATGGCCATCGT 1687
Qу
            Db
      396 CTGCCTCATGATCGGCATGGCCCACGTCCTGGTGGTCTACCGCGTCCTGGCCTCCGCGCT 455
     1688 GGTGTCCAGGTCGGGCAACACCCTTCTCGCAGCCTGGGCCTCTCGCATCGCCAGCCTCAC 1747
Qу
           456 CTTCAGCAGCTCGGCCGTGCCCTTCCTGGAGGAGCAGGTGACCACGGCCGTGGTGGTGAC 515
Db
     1748 GGGGTCTGTAGTGAACCTCGTCTTCATCCTCTCCCAAGATCTATGTATCCCTGGC 1807
Qv
           Db
       516 CGGGGCTCTGGTGCACTATGTGACCATCGTCATCATGACCAAGATCAACAGGCGCGTGGC 575
Qу
      1808 CCACGTCCTGACACGATGGGAAATGCACCGCACCCAGACCAAGTTCGAGGACGCCTTCAC 1867
          Db
      576 CCTGAAGCTTTGTGACTTCGAGATGCCCAGGACCTTCTCGGAGCGAGAGAGCAGGTTCAC 635
Qv
      1868 CCTCAAGGTGTTCATCTTCCAGTTCGTCAACTTCTACTCCTCACCCGTCTACATTGCCTT 1927
          636 CATCCGCTTCTTCACACTGCAGTTCTTCACCCATTTCTCGTCTCTCATCTACATCGCCTT 695
Db
Qv
      1928 CTTCAAGGGCAGGTTTGTGGGATACCCAGGCAACTACCACCTTGTTTGGAGTCCGCAA 1987
          696 CATCCTGGGCAGGATCAACGGCCACCCCGGGAAGTCCACGCGCCTGGCGGGCTTGTGGAA 755
Db
      1988 ---TGAGGAGTGCGCGGCTGGAGGCTGCCTGATCGAGCTGCCACAGGAGCTCCTGGTCAT 2044
QУ
             Db
      756 GCTGGAAGAGTGCCACGCCAGCGGCTGCATGATGGACCTCTTCGTGCAGATGGCCATCAT 815
      2045 CATGGTGGGCAAGCAGGTCATCAACAACATGCAGGAGGTCCTCATCCCGAAGCTAAAGGG 2104
0v
          Db
      816 CATGGGCCTGAAGCAGACGCTCAGCAACTGCGTCGAGTACCTGGTCCCGTGGGTGACCCA 875
      2105 CTGGTGGCAGAAGTTCCGGCTTCGCTCCAAGAAGAGGAAGGCGGGAGCTTCTGCAGGGGC 2164
Qv
          876 CAAGTGCC---GCTCTCTGCGGGCCTCCGAGTCCGGGCACCTGCCCCGGGACCCCGAGCT 932
Db
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Qy	2165	TAGCCAGGGCCCTGGGAGGACGACTATGAGCTTGTGCCCTGTGAGGGTCTGTTTGACGA	2224
Db	933		992
Qy	2225	GTACCTGGAAATGGTGCTGCAGTTCGGCTTCGTCACCATCTTCGTGGCCGCCTGTCCGCT	2284
Db	993	GTTCATGGAGATGATCCAGTACGGCTTCACCACCATCTTCGTGGCCGCCTTCCCGCT	1052
QУ	2285	CGCGCCGCTCTTCGCCCTGCTCAACAACTGGGTGGAGATCCGCTTGGACGCGCGCAAGTT	2344
Db	1053	$\tt GGCGCCGCTGCTCGCGCTCTTCAGCAACCTCGTGGAGGTCCGCCTGGACGCCATCAAGAT$	1112
Qy	2345	CGTCTGCGAGTACCGGCGCCCTGTGGCCGAGCGCCCCAGGACATCGGCATCTGGTTCCA	2404
Db		GGTCTGGTTGCAGCGGCCCTGGTGCCGCGCAAGGCCAAGGACATCGGGACCTGGCTGCA	
Qу		CATCCTGGCGGGCCTCACGCACCTGGCGGTCATCAGCAACGCCTTCCTCCTGGCCTTCTC	
Db		GGTGCTGGAGACCATCGGTGTGCTGGCGGTCATTGCCAATGGGATGGTCATTGCCTTCAC	
Qy Db		GTCCGACTTCCTGCCGCGCGCCTACTACCGGTGGACCCGCCCCACGACCTG	
Qy		CGCGGCTTCCTCAACTTCACGCTGGCGCGAGCCCCGTCCTCCTTCGCCGC	
Db			
Qy	2567	CGCGCACAACCGCACGTGCAGGTA	2590
Db			
Qy	2591	TCGGGCTTTCCGGGATGACGATGGACATTATTCCCAGACCTACTGGAATCTTCTTGC	2647
Db	1413		1472
Qy	2648	CATCCGCCTGGCCTTCGTCATTGTGTTTGAGCATGTGGTTTTCTCCGTTGGCCGCCTCCT	2707
Db	1473	CATCCGCCTGGCCTTCGTCATCCTCTTTGAGCACGTGGCCTTGTGCATCAAGCTCATCGC	1532
Qy	2708	GGACCTCCTGGTGCCTGACATCCCAGAGTCTGTGGAGATCAAAGTGAAGCGGGGAGTACTA	2767
Db	1533	$\tt CGCCTGGTTCGTGCCCGACATCCCTCAGTCGGTGAAGAACAAGGTTCTGGAGGTGAAGTA$	1592
Qy	2768	CCTG 2771	
Db	1593	CCAG 1596	

RESULT 13 US-09-270-767-13982

; Sequence 13982, Application US/09270767

; Patent No. 6703491

; GENERAL INFORMATION:

; APPLICANT: Homburger et al.

TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster

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; FILE REFERENCE: File Reference: 7326-094
: CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEO ID NO 13982
; LENGTH: 1282
 TYPE: DNA
 ORGANISM: Drosophila melanogaster
US-09-270-767-13982
 Ouerv Match
                   6.5%; Score 216.2; DB 3; Length 1282;
 Best Local Similarity 53.9%; Pred. No. 8.2e-36;
 Matches 496; Conservative 0; Mismatches 413; Indels 12; Gaps 2;
      1587 CGCCGCATGCTGGCCGGCTCTGTGGTGATCGTGGTGATGGTGGCCGTGGTGGTCATGTGC 1646
Qv
          169 CCCGCCACCGTGTTCAGCTTTTCAGTGGTACTGCTCCTAATTGCACTGGCCTTTGTGGCA 228
Db
      1647 CTCGTGTCTATCATCCTGTACCGTGCCATCATGGCCATCGTGGTGTCCAGGTCGGGCAAC 1706
Qv
          Db
       229 CTGCTGGCAGTGGTTGTATACCGAATGTCCATGCTGGCCGCCCTTAAAGTGGGTGCTAGT 288
      1707 ACCCTTCTCGCAGCCTGGGCCTCTCGCATCGCCAGCCTCACGGGGTCTGTAGTGAACCTC 1766
Qу
           Db
       289 CCCATGACCACCTCTAGCGCTATTGTCCTAGCCACTGCATCAGCTGCCTTTGTAAATCTG 348
      1767 GTCTTCATCCTCATCCTCTCCAAGATCTATGTATCCCTGGCCCACGTCCTGACACGATGG 1826
Qv
            349 TGCCTGCTCTATATACTTAATTATGTACAATCATTTGGCTGAGTACCTGACAGAGCTG 408
Db
      1827 GAAATGCACCGCACCCAGACCAAGTTCGAGGACGCCTTCACCCTCAAGGTGTTCATCTTC 1886
Qv
          Db
       409 GAAATGTGGCGCACTCAAACTCAGTTCGATGACTCGCTTACCCTTAAAATTTATCTGCTG 468
Qу
      1887 CAGTTCGTCAACTTCTACTCCTCACCCGTCTACATTGCCTTCTTCAAGGGCAGGTTTGTG 1946
          469 CAGTTTGTAAACTACTACGCCTCCATTTTTTACATAGCTTTCTTCAAGGGTAAATTCGTT 528
Db
      1947 GGATACCCAGGCAACTACCACACCTTGTTTGGAGTCCGCAATGAGGAGTGCGCGGCTGGA 2006
0v
          529 GGTCATCCGGGAGAGTATAATAAGCTTTTTGACTATCGGCAGGAGGAGTGCTCATCGGGT 588
Db
      2007 GGCTGCCTGATCGAGCTGGCACAGGAGCTCCTGGTCATCATGGTGGGCAAGCAGGTCATC 2066
Qv
          589 GGCTGTTTAACGGAGCTGTGCATCCAGTTAGCCATTATAATGGTTGGCAAGCAGGCATTC 648
Db
Qу
      2067 AACAACATGCAGGAGG-----TCCTCATCCCGAAGCTAAAGGGCTGGTGGCAGAAGTTC 2120
          Db
       649 AACACTATTCTTGAAGTGTATCTTCCCATGTTCTGGCGAAAGGTTTTTGGCCATTCAGGTG 708
Ov
      2121 CGGCTTCGCTCCAAGAAGAGGAAGGCGGGAGCTTCTGCAGGGGCTAGCCAGGGGCCCTGG 2180
                       709 GGCCTGTCGCGACTTTTCAACAACACCCCGAATCCAGACAAGACGAAAGACGAACGCTGG 768
Db
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2181 GAGGACGACTATGAGCTTGTGCCCTGTGAG-----GGTCTGTTTGACGAGTACCTGGAA 2234
Ov
            769 ATGCGGGATTTCAAGCTACTGGATTGGGGTGCCCGAGGTCTGTTTCCCGAGTATTTGGAG 828
Db
      2235 ATGGTGCTGCAGTTCGGCTCACCATCTTCGTGGCCGCCTGTCCGCTCGCCCGCTC 2294
Qv
           Db
      829 ATGGTCTTGCAGTACGGCTTCGTAACCATCTTTGTGGCCGCTTTTCCGCTGGCGCCATTC 888
     2295 TTCGCCCTGCTCAACAACTGGGTGGAGATCCGCTTGGACGCGCGCAAGTTCGTCTGCGAG 2354
Qy
          Db
      889 TTTGCCCTGCTAAATAATATCTTGGAAATGCGACTGGATGCAAAGAAACTATTGACCCAC 948
     2355 TACCGGCGCCCTGTGGCCGAGCGCCCCAGGACATCGGCATCTGGTTCCACATCCTGGCG 2414
Qу
           Db
      949 CACAAGCGTCCAGTATCACAGCGAGTTCGAGATATAGGAGTGTGGTATCGTATCCTGGAC 1008
     2415 GGCCTCACGCACCTGGCGGTCATCAGCAACGCCTTCCTCCTGGCCTTCTCGTCCGACTTC 2474
Qv
           Db
      1009 TGCATAGGCAAGCTCAGCGTGATCACAAATGGATTCATCATAGCCTTTACCTCTGACATG 1068
Qv
     2475 CTGCCGCGCGCCTACTACCGG 2495
           1 11111 111111
Db
     1069 ATTCCGCGTTTGGTGTACCGG 1089
RESULT 14
US-10-104-047-1146
; Sequence 1146, Application US/10104047
; Patent No. 6943241
; GENERAL INFORMATION:
; APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: No. 6943241el full length cDNA
; FILE REFERENCE: H1-A0105
; CURRENT APPLICATION NUMBER: US/10/104,047
; CURRENT FILING DATE: 2002-03-25
; PRIOR APPLICATION NUMBER:
; PRIOR FILING DATE:
; NUMBER OF SEQ ID NOS: 4096
: SOFTWARE: Patentin Ver. 2.1
; SEO ID NO 1146
; LENGTH: 2293
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-104-047-1146
 Query Match
                   5.2%; Score 170.8; DB 3; Length 2293;
 Best Local Similarity 54.9%; Pred. No. 4.1e-26;
 Matches 400; Conservative 0; Mismatches 322; Indels 6; Gaps 3;
Oy 1759 TGAACCTCGTCTTCATCCTCATCCTCCAAGATCTATGTATCCCTGGCCCACGTCCTGA 1818
           324 TCAACTTCTGTATCATTATGTTGCTGAATGTGCTCTATGAAAAAGTTGCCCTGCTTCTGA 383
Db
Qy 1819 CACGATGGGAAATGCACCGCACCCAGACCAAGTTCGAGGACGCCTTCACCCTCAAGGTGT 1878
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384 CGAATTTAGAACAGCCTCGCACAGAGTCTGAGTGGGAGAACAGCTTCACCCTGAAAATGT 443

Db

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1879 TCATCTTCCAGTTCGTCAACTTCTACTCCTCACCCGTCTACATTGCCTTCTTCAAGGGCA 1938
Qv
          11.1
Db
       444 TTCTTTTCAGTTTGTCAATCTGAACAGCTCCACATTTTACATCGCATTCTTCCTCGGAA 503
      1939 GGTTTGTGGGATACCCAGGCAACTACCACACCTTGTT---TGGAGTCCGCAATGAGGAGT 1995
Qy
          504 GATTTACAGGACACCCAGGTGCCTACTTGAGGCTGATAAACAGGTGGAGACTAGAAGAGT 563
Db
      1996 GCGCGGCTGGAGCTGCCTGATCGAGCTGCACAGGAGCTCCTGGTCATCATGGTGGGCA 2055
QУ
          Dh
       564 GCCACCCTAGTGGATGCCTTATTGATCTGTGTATGCAAATGGGTATTATAATGGTGCTAA 623
      2056 AGCAGGTCATCAACAACATGCAGGAGGTCCTCATCCCGAAGCTAAAGGGCTGGTGGCAGA 2115
Qv
          624 AGCAGACCTGGAATAATTTCATGGAACTTGGCTACCCGTTAATTCAGAATTGGTGGACTA 683
Db
     2116 AGTTCCGGCTTCGCTCCAAGAAGAGGGAGGCGGGGGCTTCTGCAGGGGCTAGCCAGGGGC 2175
Qy
                684 GAAGAAAGTACG--ACAAGAACATGGACCTGAAAGGAAAATAAGTTTCCCACAATGGG- 740
Db
     2176 CCTGGGAGGACGACTATGAGCTTGTGCCCTGTGAGGGTCTGTTTGACGAGTACCTGGAAA 2235
Qу
             Db
      741 AAAAGGACTATAACCTTCAGCCGATGAATGCCTATGGACTCTTCGATGAATACTTAGAAA 800
      2236 TGGTGCTGCAGTTCGGCTCACCATCTTCGTGGCCGCCTGTCCGCTCGCGCCGCTCT 2295
Qv
          801 TGATTCTTCAGTTTGGATTCACAACTATCTTTGTGGCAGCTTTTCCCCTAGCACCACTTC 860
Db
     2296 TCGCCCTGCTCAACAACTGGGTGGAGATCCGCTTGGACGCGCGCAAGTTCGTCTGCGAGT 2355
Qy
          Db
       861 TGGCCTTACTGAATAACATAATTGAAATTCGACTTGATGCTTACAAATTTGTCACACAGT 920
Qу
      2356 ACCGGCGCCTGTGGCCGAGCGCCCAGGACATCGGCATCTGGTTCCACATCCTGGCGG 2415
            Db
      921 GGAGGAGACCTTTAGCTTCAAGGGCCAAAGACATAGGAATTTGGTATGGAATTCTTGAAG 980
Qv
      2416 GCCTCACGCACCTGGCGGTCATCAGCAACGCCTTCCTCCTGGCCTTCTCGTCCGACTTCC 2475
                 Db
      981 GCATTGGAATTCTCTCTGTTATCACAAATGCATTTGTCATAGCGATAACATCTGACTTTA 1040
Qv
      2476 TGCCGCGC 2483
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Db
     1041 TCCCTCGC 1048
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RESULT 15 US-10-100-683-1599

[;] Sequence 1599, Application US/10100683; Patent No. 7368531

[;] GENERAL INFORMATION:

[;] APPLICANT: Rosen, et al.

[;] TITLE OF INVENTION: Human Secreted Proteins

[;] FILE REFERENCE: PS900

[;] CURRENT APPLICATION NUMBER: US/10/100,683

[;] CURRENT FILING DATE: 2002-03-19

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PRIOR APPLICATION NUMBER: US 60/040,162
; PRIOR FILING DATE: 1997-03-07
; PRIOR APPLICATION NUMBER: US 60/043,576
; PRIOR FILING DATE: 1997-04-11
; PRIOR APPLICATION NUMBER: US 60/047,601
; PRIOR FILING DATE: 1997-05-23
; PRIOR APPLICATION NUMBER: US 60/056,845
; PRIOR FILING DATE: 1997-08-22
; PRIOR APPLICATION NUMBER: US 60/043,580
  PRIOR FILING DATE: 1997-04-11
; PRIOR APPLICATION NUMBER: US 60/047,599
  PRIOR FILING DATE: 1997-05-23
 PRIOR APPLICATION NUMBER: US 60/056,664
  PRIOR FILING DATE: 1997-08-22
 PRIOR APPLICATION NUMBER: US 60/043,314
  PRIOR FILING DATE: 1997-04-11
 PRIOR APPLICATION NUMBER: US 60/047,632
; PRIOR FILING DATE: 1997-05-23
; PRIOR APPLICATION NUMBER: US 60/056,892
 PRIOR FILING DATE: 1997-08-22
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEO ID NOS: 13468
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1599
; LENGTH: 2371
 TYPE: DNA
  ORGANISM: Homo sapiens
US-10-100-683-1599
 Query Match
                     4.8%; Score 157.2; DB 7; Length 2371;
 Best Local Similarity 61.9%; Pred. No. 3.2e-23;
 Matches 249; Conservative 0; Mismatches 153; Indels 0; Gaps 0;
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      2181 GAGGACGACTATGAGCTTGTGCCCTGTGAGGGTCTGTTTGACGAGTACCTGGAAATGGTG 2240
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          6 GAGGTGGATTACAACCTGGAGCCCTTCGCGGGCCTCACCCCAGAGTACATGGAAATGATC 65
Qv
      2241 CTGCAGTTCGGCTTCGTCACCATCTTCGTGGCCGCCTGTCCGCTCGCCGCCGCTCTTCGCC 2300
            66 ATCCAGTTTGGCTTCGTCACCCTGTTTGTCGCCTCCTTCCCCCTGGCCCCACTGTTTGCG 125
Db
Qv
       2301 CTGCTCAACAACTGGGTGGAGATCCGCTTGGACGCGCGCAAGTTCGTCTGCGAGTACCGG 2360
           126 CTGCTGAACAACATCATCGAGATCCGCCTGGACGCCAAAAAGTTTGTCACTGAGCTCCGA 185
Db
       2361 CGCCCTGTGGCCGAGCGCCCAGGACATCGGCATCTGGTTCCACATCCTGGCGGGCCTC 2420
Qу
            Db
       186 AGGCCGGTAGCTGTCAGAGCCAAAGACATCGGAATCTGGTACAATATCCTCAGAGGCATT 245
       2421 ACGCACCTGGCGGTCATCAGCAACGCCTTCCTCCTGGCCTTCTCGTCCGACTTCCTGCCG 2480
0v
             Db
        246 GGGAAGCTTGCTGTCATCATCATGCCTTCGTGATCTCCTTCACGTCTGACTTCATCCCG 305
       2481 CGCGCCTACTACCGGTGGACCCGCGCCCACGACCTGCGCGGCTTCCTCAACTTCACGCTG 2540
Qv
                 306 CGCCTGGTGTACCTCTACATGTACAGTAAGAACGGGACCATGCACGGCTTCGTCAACCAC 365
Db
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QУ	2541 GCGCGAGCCCCGTCCTTCGCCGCCGCGCACAACCGCACG 2582
Db	366 ACCCTCCCCCCTCCAACGTCAGTGACTTCCAGAACGGCACG 407

Search completed: March 16, 2009, 16:47:17 Job time : 1176 secs